

2879099\_1.TXT  
SEQUENCE LISTING

<110> Horiguchi, Takashi  
Matsui, Hideki  
Watanabe, Tomomichi

<120> NOVEL PROTEIN AND ITS DNA

<130> Q101074

<140> US 10/547,843

<141> 2005-09-06

<150> PCT/JP2004/003033

<151> 2004-03-09

<150> JP2003-64197

<151> 2003-03-10

<150> JP2003-149679

<151> 2003-05-27

<160> 10

<210> 1

<211> 264

<212> PRT

<213> Homo sapiens

<400> 1

```

Met Gly Gly Ala Gln Leu Glu Leu Pro Ser Gly Ala Arg Pro Gly Val
 1           5           10           15
Cys Val Arg Arg Ser Phe Arg Ala His Ala Gly Asp Gln Pro Arg Arg
      20           25           30
Pro Pro Gly Pro Ile Pro Val Pro Gly Thr Met Lys Gln Glu Ser Ala
      35           40           45
Ala Pro Asn Thr Pro Pro Thr Ser Gln Ser Pro Thr Pro Ser Ala Gln
      50           55           60
Phe Pro Arg Asn Asp Gly Asp Pro Gln Ala Leu Trp Ile Phe Gly Tyr
 65           70           75           80
Gly Ser Leu Val Trp Arg Pro Asp Phe Ala Tyr Ser Asp Ser Arg Val
      85           90           95
Gly Phe Val Arg Gly Tyr Ser Arg Arg Phe Trp Gln Gly Asp Thr Phe
      100          105          110
His Arg Gly Ser Asp Lys Met Pro Gly Arg Val Val Thr Leu Leu Glu
      115          120          125
Asp His Glu Gly Cys Thr Trp Gly Val Ala Tyr Gln Val Gln Gly Glu
      130          135          140
Gln Val Ser Lys Ala Leu Lys Tyr Leu Asn Val Arg Glu Ala Val Leu
 145          150          155          160

```

# 2879099\_1.TXT

Gly Gly Tyr Asp Thr Lys Glu Val Thr Phe Tyr Pro Gln Asp Ala Pro  
165 170 175  
Asp Gln Pro Leu Lys Ala Leu Ala Tyr Val Ala Thr Pro Gln Asn Pro  
180 185 190  
Gly Tyr Leu Gly Pro Ala Pro Glu Glu Ala Ile Ala Thr Gln Ile Leu  
195 200 205  
Ala Cys Arg Gly Phe Ser Gly His Asn Leu Glu Tyr Leu Leu Arg Leu  
210 215 220  
Ala Asp Phe Met Gln Leu Cys Gly Pro Gln Ala Gln Asp Glu His Leu  
225 230 235 240  
Ala Ala Ile Val Asp Ala Val Gly Thr Met Leu Pro Cys Phe Cys Pro  
245 250 255  
Thr Glu Gln Ala Leu Ala Leu Val  
260

<210> 2  
<211> 792  
<212> DNA  
<213> Homo sapiens

<400> 2  
atggggggcg ctcagctgga gctaccgagc ggtgccaggc caggtgtgtg cgtccgtcgg 60  
tctttccgtg cccacgccgg agaccagccc cggaggccgc ctgggcctat ccctgtgcc 120  
ggcaccatga agcaggagtc tgcagccccc aacaccccgc ccacctcgca gtcacctacg 180  
ccgtccgtc agttccccg aaacgacggc gacctcaag cgctgtggat ttctgggtac 240  
ggctccctgg tgtggaggcc cgacttcgcc tacagcgaca gccgtgtggg cttcgtgcgc 300  
ggctacagcc gccgtttctg gcaggagac accttccatc ggggcagcga caagatgcct 360  
ggccgtgtgg tgacgctcct tgaagatcat gagggctgca cttggggcgt ggcataccaa 420  
gtgcaagggg agcaggtaag caaggccctg aagtaccta atgtgcgaga ggcagtgtt 480  
ggtggctacg ataccaagga ggtcaccttc tatccccaag atgtcctga ccaaccactg 540  
aaggcattgg cctatgtggc caccacacag aacctggtt acctgggccc tgcgcctgaa 600  
gaggccattg ccacgcagat cctggcctgc cggggcttct ccggccacaa cttgaatac 660  
ttgctgcgtc tggcagactt catgcagctc tgtgggcctc aggcgcagga cgagcacctg 720  
gcagccatcg tggacgctgt gggcaccatg ttgcctgct tctgcccac cgagcaggct 780  
ctggcgctgg tg 792

<210> 3  
<211> 24  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Primer

<400> 3	
gcaggcaggg gtataagaca caaa	24
<210> 4	
<211> 24	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 4	
gccacccgcc atgccttctc agct	24
<210> 5	
<211> 24	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 5	
cgccagggtt ttcccagtca cgac	24
<210> 6	
<211> 27	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 6	
agcggataac aatttcacac aggaaac	27
<210> 7	
<211> 30	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 7	
gccatggggg gcgctcagct ggagctaccg	30
<210> 8	
<211> 26	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Primer	
<400> 8	
tcacaccagc gccagagcct gctcgg	26
<210> 9	
<211> 20	
<212> DNA	
<213> Artificial Sequence	

# 2879099\_1.TXT

<220>		
<223>	Primer	
<400>	9	
	taatacgact cactataggg	20
<210>	10	
<211>	18	
<212>	DNA	
<213>	Artificial Sequence	
<220>		
<223>	Primer	
<400>	10	
	tagaaggcac agtcgagg	18

PAGE: 1  
12/10/2007

VERIFICATION SUMMARY REPORT

DATE:

PATENT APPLICATION

TIME:

14:18:07

INPUT SEQ: H:\PatIn31\Q101074revSeqList.TXT

GENERAL INFORMATION SECTION

-----  
4,<110> Horiguchi, Takashi  
5, MATSUI, Hideki  
6, WATANABE, Tomomichi  
8,<120> NOVEL PROTEIN AND ITS DNA  
10,<130> Q101074  
12,<140> US 10/547,843  
14,<141> 2005-09-06  
16,<150> PCT/JP2004/003033  
17,<151> 2004-03-09  
19,<150> JP2003-64197  
20,<151> 2003-03-10  
22,<150> JP2003-149679  
23,<151> 2003-05-27  
25,<160> 10

ERRORED LINES SECTION

-----  
STATISTICS SUMMARY  
-----

Application Serial Number: US 10/547,843  
Alpha or Numeric or Xml: Numeric  
Application Class:  
Application File Date: 2005-09-06  
Art Unit:  
Software Application:  
Total Number of Sequences: 10  
Total Nucleotides: 985  
Total Amino Acids: 264  
Number of Errors: 0  
Number of Warnings: 0  
Number of Corrections: 0